

PATENT
3645-0104P

IN THE U.S. PATENT AND TRADEMARK OFFICE

Applicant: Richard B. THOMPSON et al. Conf.: 9931
Appl. No.: 09/942,708 Group:
Filed: August 31, 2001 Examiner:

For: DETERMINATION OF METAL IONS IN SOLUTION
BY PHOTOLUMINESCENCE ANISOTROPY

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LETTER TO THE OFFICIAL DRAFTSPERSON TC 170

Assistant Commissioner for Patents
Washington, DC 20231

December 11, 2001

Sir:

Attached hereto is/are one (1) sheet(s) of formal drawings which comply with the provisions of 37 C.F.R. § 1.84. The drawings should be made a part of the record of the above-identified application.

If necessary, the Commissioner is hereby authorized in this, concurrent, and future replies, to charge payment or credit any overpayment to Deposit Account No. 02-2448 for any additional fees required under 37 C.F.R. §§1.16 or 1.17; particularly, extension of time fees.

Respectfully submitted,

BIRCH, STEWART, KOLASCH & BIRCH, LLP

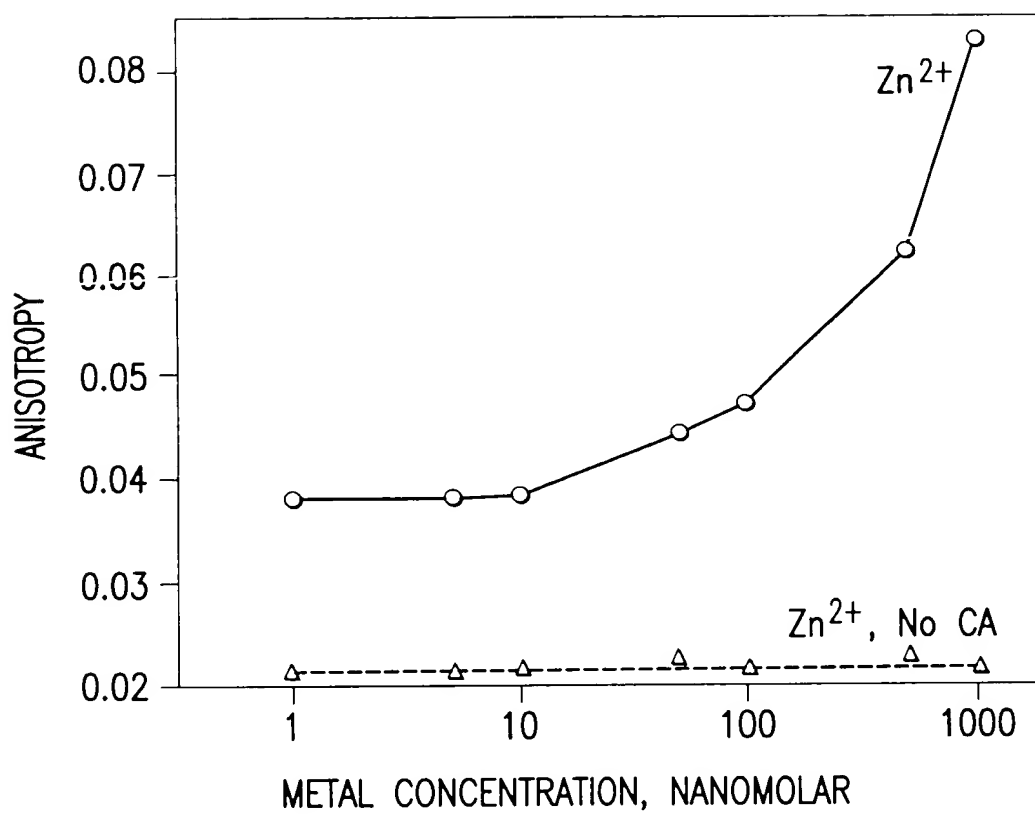
By Mark J. Nuell, #36,623

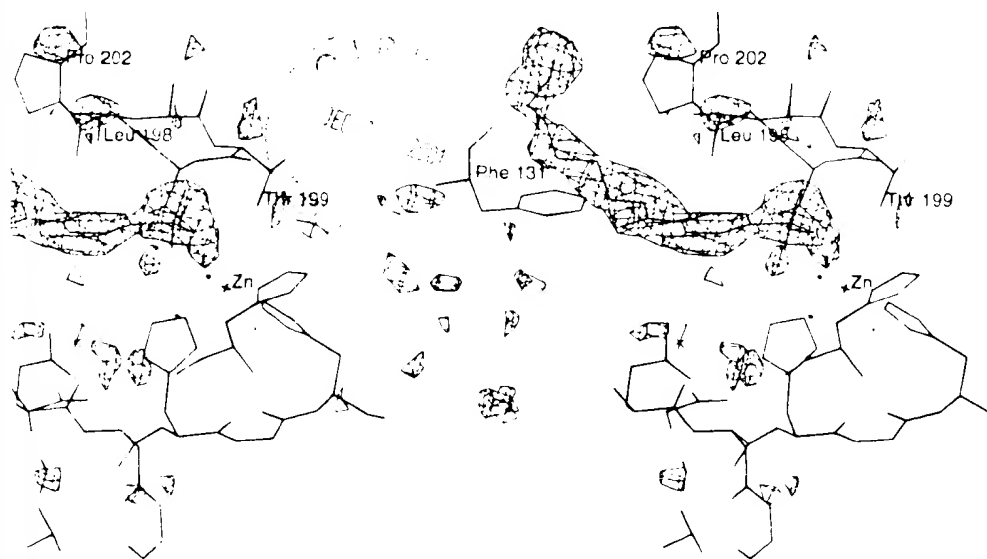
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Attachments

Figure 9





CAII-**3** complex contoured at 2.4σ . Enzyme residues Phe-131, Leu-198, Thr-199, Pro-202, and Zn^{2+} are shown in contact with the thiocarbonyl moiety. No electron density is observed for the fluorescein portion of compound **3**.

ns

separation (Å)	
2	3
2.1	2.4
2.8 ^a	3.1 ^a
3.1 ^a	3.5
3.3	3.3
3.3	3.6
2.9 ^a	
3.3	2.9
3.0 ^a	

deduced from distance and

hydrogen bond contacts with the enzyme. Thus, only hydrophobic interactions appear to stabilize the binding of **2**. The protein residues involved in binding **2**. This is consistent with the extensive hydrophobic interactions in the 198 region stabilize the binding of **2** in a nearly identical

As shown above, there are two hydrogen bonds between CAII and **2**. The hydrogen bond from water 401, and the hydrogen bond from the side chain of the active site residue to water 302,

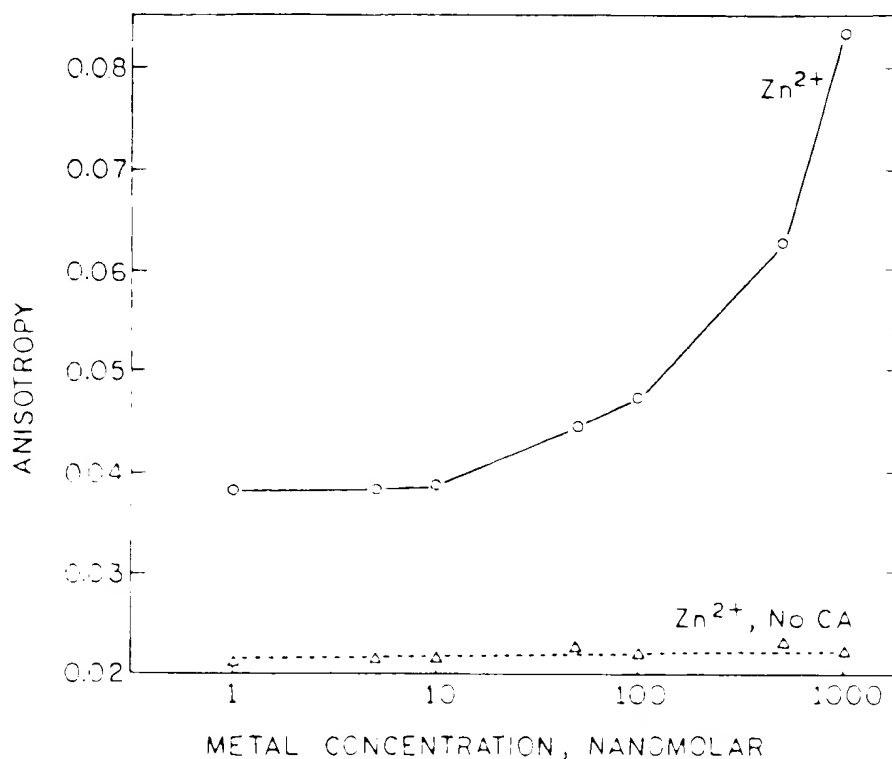


Figure 5. Zn^{2+} concentration-dependent fluorescence anisotropies of $1 \mu\text{M}$ **3** are depicted in the absence (triangles) and presence (circles) of $1 \mu\text{M}$ apo-CAII. Anisotropy = $(I_{\parallel} - I_{\perp}) / (I_{\parallel} + 2I_{\perp})$; see eq 6.

An electron density map of the CAII-**3** complex is shown in Figure 4, and selected enzyme-inhibitor interactions are recorded in Table 2. We note that the hydrophobic and hydrogen bond interactions described in the previous paragraph are presumably sufficient to stabilize inhibitor binding to the apoenzyme, although with $\sim 10^3$ -fold weaker affinity, in the absence of sulfonamide zinc coordination.